Figure 1

```
ggcgcccgcc cgccgcccgc ccggcccccg gctgcctccc ttcctccctc cctctctttc tcccttgcgc tcgctcgctc gctcgccctc ggcgcatggg      100


ccccgcgccg ggccccgggg cctcgggccg cctggcctcc ggggtccccct aggcccgggc gtgggcgggg cagcccggcc tgacgcagcc tctgtacccc      200


accaccacca ccaccagggc cggcggcggc ggctgccccg agggacgggg ccctaggcgg tggcgATGGG GGCCGTCCGG ATCGCGCCCG GCCTGGCGCT      300
                                                                         M  G  A  V  R  I  A  P  G  L  A  L

GCTGCTCTGC TGCCCGGTGC TCAGCTCCGC GTACGCGCTG GTGGATGCAG ATGACGTCAT GACCAAAGAG GAGCAGATCT TCCTGCTGCA CCGCGCCCAG      400
 L  L  C  C  P  U  L  S  S  A  Y  A  L  V  D  A  D  D  V  M  T  K  E  E  Q  I  F  L  L  H  R  A  Q

GCCCAGTGCC AGAAGCGGCT CAAAGAAGTC CTGCAGAGGC CAGCTGACAT AATGGAATCA GACAAAGGAT GGGCTTCTGC ATCCACATCA GGGAAGCCTA      500
 A  Q  C  Q  K  R  L  K  E  U  L  Q  R  P  A  D  I  M  E  S  D  K  G  W  A  S  A  S  T  S  G  K  P  K

AGAAAGAGAA GGCATCTGGG AAGCTCTACC CTGAGTCCGA GGAGGACAAG GAGGTGCCCA CTGGCAGCAG GCACCGAGGG CGCCCCTGCC TGCCCGAGTG      600
 K  E  K  A  S  G  K  L  Y  P  E  S  E  E  D  K  E  U  P  T  G  S  R  H  R  G  R  P  C  L  P  E  W

GGACCACATC CTTTGCTGGC CGCTGGGGGC ACCAGGTGAG GTGGTGGCTG TGCCCTGTCC CGACTACATT TATGACTTCA ATCACAAAGG CCATGCCTAC      700
 D  H  I  L  C  W  P  L  G  A  P  G  E  U  U  A  U  P  C  P  D  Y  I  Y  D  F  N  H  K  G  H  A  Y

CGTCGCTGTG ACCGCAATGG CAGCTGGGAG CTGGTGCCTG GACACAACCG GACGTGGGCC AACTACAGCG AGTGTGTCAA GTTCCTGACC AACGAGACTC      800
 R  R  C  D  R  N  G  S  W  E  L  U  P  G  H  N  R  T  W  A  N  Y  S  E  C  U  K  F  L  T  N  E  T  R
                                                                     I
GTGAACGGGA GGTGTTTGAC CGC[CTGGGGCA TGATCTACAC CGTGGGCTAC TCCGTGTCGC TGGCCTCCCT CACCGTGGCC GTGCTCATCC TGGCC]TACTT      900
 E  R  E  U  F  D  R  L  G  M  I  Y  T  U  G  Y  S  U  S  L  A  S  L  T  U  A  U  L  I  L  A  Y  F
                                                                         II
CAGGCGGCTG CACTGCACAC GCAACTACAT CCAC[ATGCAC CTGTTCCTGT CCTTCATGCT TCGCGCCGTG AGCATCTTCG TCAAGGACGC GGTGCTCTAC     1000
 R  R  L  H  C  T  R  N  Y  I  H  M  H  L  F  L  S  F  M  L  R  A  U  S  I  F  U  K  D  A  U  L  Y
 _____
[TCGGGCGCC]A CGCTCGACGA GGCCGAGCGC CTCACGGAGG AAGAGCTGCG CGC[CATGGCC CAGGCACCCC CGCCGCCCAC CGCCGCCGCC GGCTACGCGG     1100
 S  G  A  T  L  D  E  A  E  R  L  T  E  E  E  L  R  A  I  A  Q  A  P  P  P  P  T  A  A  G  Y  A  G
                                                                             III
GCTGCAGGGT AGCTGTGACC TTCTTCCTTT ATTTCCTGGC CACCAACTAC TAC[TGGATTC TGGTGGAGGG GCTGTACCTG CATAGTCTCA TCTTCATGGC     1200
 C  R  U  A  U  T  F  F  L  Y  F  L  A  T  N  Y  Y  W  I  L  U  E  G  L  Y  L  H  S  L  I  F  M  A
                                                                     IV
[CTTCTTC]TCA GAGAAGAAGT ACCTG[TGGGG CTTCACGGTC TTCGGCTGGG GTCTGCCCGC CGTCTTCGTG GCTGTGTGGG TCAGCGTGAG AGCCACCCTG     1300
 F  F  S  E  K  K  Y  L  W  G  F  T  U  F  G  W  G  L  P  A  U  F  U  A  U  W  U  S  U  R  A  T  L
                                                                             V
[GCCAACACG]G GGTGCTGGGA CTTGAGCTCC GGGAACAAGA AGTGGATC[AT CCAGGTGCCC ATCCTGGCCT CTATTGTGCT CAACTTCATC TTGTTCATCA     1400
 A  N  T  G  C  W  D  L  S  S  G  N  K  K  W  I  I  Q  U  P  I  L  A  S  I  U  L  N  F  I  L  F  I

[ACATCGTCCG GGTGCTCGCC ACA]AAGCTGC GGGAGACCAA TGCCGGCCGG TGTGACACGC GGCAGCAGTA CCGGAAG[CTG CTCAAATCCA CACTGGTGCT     1500
 I  U  R  U  L  A  T  K  L  R  E  T  N  A  G  R  C  D  T  R  Q  Q  Y  R  K  L  L  K  S  T  L  U  L
    VI
[CATGCCGCTC TTTGGCGTCC ACTACATCGT CTTCATGGCC ACGCCG]TACA CCGAGGTCTC AGGGACGCTC TGGCAAGTCC AGATGCACTA CGAG[ATGCTG     1600
 M  P  L  F  G  U  H  Y  I  U  F  M  A  T  P  Y  T  E  U  S  G  T  L  W  Q  U  Q  M  H  Y  E  M  L
            VII
[TTCAACTCCT TCCAGGGATT TTTTGTCGCC ATCATATACT GTTTCTGCAA T]GGCGAGGTA CAGGCCGAGA TCAAGAAATC CTGGAGCCGC TGGACACTGG     1700
 F  N  S  F  Q  G  F  F  U  A  I  I  Y  C  F  C  N  G  E  U  Q  A  E  I  K  K  S  W  S  R  W  T  L  A

CCCTGGACTT CAAGCGCAAG GCCCGAAGTG GGAGCAGCAG TTACAGCTAC GGCCCCGATGG TGTCTCACAC GAGCGTGACC AACGTAGGCC CCCGCGCGGG     1800
 L  D  F  K  R  K  A  R  S  G  S  S  S  Y  S  Y  G  P  M  U  S  H  T  S  U  T  N  U  G  P  R  A  G

ACTTGGCCTG CCCCTCAGCC CCCGCCTGCT GCCCGCCGCT GCCGCCACCA CCACCGCCAC CACCAACGGC CACCCCCCGA TCCCGGGCCA CACCAAGCCA     1900
 L  G  L  P  L  S  P  R  L  L  P  A  A  A  A  T  T  T  A  T  T  N  G  H  P  P  I  P  G  H  T  K  P

GGGGCCCCGA CCCTCCCGGC CACACCACCT GCCACGGCTG CTCCCAAGGA CGATGGGTTC CTCAACGGCT CCTGCTCGGG GCTGGACGAG GAGGCCTCCG     2000
 G  A  P  T  L  P  A  T  P  P  A  T  A  A  P  K  D  D  G  F  L  N  G  S  C  S  G  L  D  E  E  A  S  A

CGCCGGAGCG GCCTCCCGCC CTGCTGCAGG AGGAGTGGGA GACGGTCATG TGAtcgggga cctgtgccag ggttggactc gtggacataa gggccgacag     2100
 P  E  R  P  P  A  L  L  Q  E  E  W  E  T  U  M  .


acggaccaag agacaggcgg ttggacagtt gcccactcag ggctggggct gggaagacaa aacaaaaaaa aaaaaaa                                2177
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SEQ. ID No. 1

Figure 2

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dPTH1  ATGGGGGCCG TCCGGATCGC GCCCGGCTG GCGCTGCTGC TCTGCTGCCC GGTGCTCAGC TCCGCGTACG CGCTG  75
rPTH1  ATGGGGGCCG CCCGGATCGC ACCCGGCCTG GCGCTGCTAC TCTGCTGCCC AGTGCTCAGC TCCGCATATG CGCTG  75
mPTH1  ATGGGGGACCG CCCGGATCGC ACCCGGCCTG GCGCTGCTTC TCTGCTGCCC AGTGCTCAGC TCCGCATATG CGCTG  75
hPTH1  ATGGGGGACCG CCCGGATCGC ACCCGGCCTG GCGCTGCTGC TCTGCTGCCC CGTGCTCAGC TCCGCGTACG CGCTG  75

dPTH1  GTGGATGCAG ATGACGTCAT GACCAAAGAG GAACAGATCT TCCTGCTGCA CCGCGGCCAG GCCCAGTGCC AGAAG  150
rPTH1  GTGGACGCAG ACGATGTCTT TACCAAAGAG GAACAGATTT TCCTGCTGCA CCGTGCCCAG GCGCAATGTG ACAAG  150
mPTH1  GTGGACGCAG ACGATGTCTT TACCAAAGAG GAACAGATTT TCCTGCTGCA CCGTGCCCAG GCGCAATGTG ACAAG  150
hPTH1  GTGGATGCAG ATGACGTCAT GACTAAAGAG GAACAGATCT TCCTGCTGCA CCGTGCTCAG GCCCAGTGCG AAAAA  150

dPTH1  CCGCTCAAGG AAGTCCTGCA GAGGCCAGCT GACATAATGG AATCAGACAA AGGATGGGCT TCTGCATCCA CATCA  225
rPTH1  CTTGCTCAAGG AAGTTCTGCA CACAGCAGCC AACATAATGG AGTCAGACAA GGGGTGGACA CCAGCATCTA CGTCA  225
mPTH1  CTTGCTCAAGG AAGTTCTGCA CACAGCAGCC AGCATAATGG AATCAGACAA GGGGATGGACA TCTGCGTCA CATCA  225
hPTH1  CCGCTCAAGG AAGGTCCTGCA GAGGCCAGCC AGCATAATGG AATCAGACAA GGGGATGGACA TCTGCGTCA CATCA  225

dPTH1  GGGAAGCCTA AGAAAGAGAA GGCATCTGGG AAGTCTACC CTGAGTCCGA AGAGGACAAG GAGGTGCCCA CTGGC  300
rPTH1  GGGAAGCCTA GGAAAGAGAA GGCATCGGGA AAGTTCTACC CTGAGTCTAA AGAGAACAAG GATGTGCCCA CGGGC  300
mPTH1  GGGAAGCCTA GGAAAGAGAA GGCACGGGA AAGTTCTACC CGGAGTCTAA AGAGAACAAG GATGTGCCCA CGGGC  300
hPTH1  GGGAAGCCTA GGAAAGATAA GGCATCTGGG AAGTCTACC CTGAGTCTGA GGAGGACAAG GAGGCACCCA CTGGC  300

dPTH1  AGCAGGCACC GAGGGCGCCC CTGCCTGCCC GAGTGGGACC ACATCCTTTG CTGGCCGCTG GGGGCACCAG GTGAG  375
rPTH1  AGCAGGCGCA GAGGGCGTCC CTGTCTGCCC GAGTGGGACA ACATCGTTTG CTGGCCATTA GGGGCACCAG GTGAA  375
mPTH1  AGCAGGCGCC GAGGGCGTCC CTGTCTGCCA GAGTGGGACA ACATCGTTTG CTGGCCATTG GGGGCACCAG GTGAA  375
hPTH1  AGCAGGTACC GAGGGCGCCC CTGTCTGCCG GAATGGGACC ACATCCTCTG CTGGCCGCTG GGGGCACCAG GTGAG  375

dPTH1  GTGGTGGCTG TGCCCTGTCC CGACTACATT TATGACTTCA ATCACAAAGG CCATGCCTAC CGTCGCTGTG ACCGC  450
rPTH1  GTGGTGGCAG TACCTTGTCC CGATTACATT TATGACTTCA ATCACAAAGG CCATGCCTAC AGACGCTGTG ACCGC  450
mPTH1  GTGGTGGCAG TACCTTGTCC CGATTACATT TATGACTTCA ACCACAAAGG CCATGCCTAC AGACGCTGCG ACCGC  450
hPTH1  GTGGTGGCTG TGCCCTGTCC GGACTACATT TATGACTTCA ATCACAAAGG CCATGCCTAC CGACGCTGTG ACCGC  450

dPTH1  AATGGCAGCT GGGAGCTGGT GCCTGGACAC AACCGGACGT GGGCCAACTA CAGCGAGTGT GTCAAGTTCC TGACC  525
rPTH1  AATGGCAGCT GGGAGCTGGT TCCAGGGCAC AACCGGACGT GGGCCAACTA CAGCGAGTGC CTCAAGTTCA TGACC  525
mPTH1  AATGGCAGCT GGGAGCTGGT TCCAGGGCAC AACCGGACGT GGGCCAACTA CAGCGAGTGT CTCAAGTTTC TGACC  525
hPTH1  AATGGCAGCT GGGAGCTGGT GCCTGGACAC AACAGGACGT GGGCCAACTA CAGCGAGTGT GTCAAGTTCC TGACC  525

dPTH1  AACGAGACTC GTGAACGGGA GGTGTTTGAC CGCCTGGGCA TGATCTACAC CGTGGGCTAC TCCGTGTCGC TGGCC  600
rPTH1  AATGAGACGC GGGAACGGGA GGTATTTGAC CGCCTGGGCA TGATCTACAC CGTGGGATAT TCCATGTCTC TTGCC  600
mPTH1  AATGAGACTC GGGAACGGGA GGTATTTGAC CGCCTGGGCA TGATTTACAC CGTGGGCTAC TCCGTGTCTC TTGCG  600
hPTH1  AATGAGACTC GTGAACGGGA GGTGTTTGAC CGCCTGGGCA TGATTTACAC CGTGGGCTAC TCCGTGTCGC TGGCG  600

dPTH1  TCCCTCACGG TGGCGGTGCT CATCCTGGCC TACTTCAGGC GGCTGCACTG CACACGCAAC TACATCCACA TGCAC  675
rPTH1  TCCCTCACGG TGGCTGTGCT CATCCTGGCC TATTTTAGGC GGCTGCACTG CACGGCGCAAC TACATCCACA TGCAC  675
mPTH1  TCCCTCACGG TGGCTGTGCT CATCCTAGCC TATTTTAGGC GGCTGCACTG CACGGCGCAAC TACATCCACA TGCAC  675
hPTH1  TCCCTCACGG TAGCTGTGCT CATCCTAGCC TACTTTAGGC GGCTGCACTG CACGGCGCAAC TACATCCACA TGCAC  675

dPTH1  CTGTTCCTGT CCTTCATGCT TCGCGCCGTG AGCATCTTCG TCAAGGACGC GGTGCTCTAC TCGGGCGGCA CGCTC  750
rPTH1  ATGTTCCTGT CGTTTATGCT GCGCGCCGGG AGCATCTTCG TCAAGGACGC TGTGCTCTAC TCTGGCTTCA CGCTG  750
mPTH1  ATGTTCCTGT CCTTTATGCT GCGCGCCGTG AGCATCTTCG TCAAGGACGC TGTGCTCTAC TCTGGCTTCA CGCTG  750
hPTH1  CTGTTCCTGT CCTTCATGCT GCGCGCCGTG AGCATCTTCG TCAAGGACGC TGTGCTCTAC TCTGGCGGCA CGCTT  750

dPTH1  GACGAGGCTG AGCGCCTCAC GGAGGAAGAG CTGCGCGCA TCGCCCAGGC ACCCCCGCCG CCCACGCCGG CCGCC  825
rPTH1  GATGAGGCTG AGCGCCTCAC GGAGGAAGAG TTGCACATCA TCGCGCAGG- ---------- --TGCCACCT CCGCC  812
mPTH1  GATGAGGCTG AGCGCCTCAC GGAGGAAGAG TTGCATATCA TCGCGCAGG- ---------- --TGCCGCCT CCGCC  812
hPTH1  GATGAGGCTG AGCGCCTCAC CGAGGAAGAG CTGCGCGCA TCGCCCAGGC GCCCCCGCCG CCTGCCACCG CCGC-  824

dPTH1  GGC------ --------- T ACGCGGGCTG CAGGGTAGCT GTGACCTTCT TCCTTTAGTT CCTGGCTACC AACTA  884
rPTH1  GGCCGCTGCC GCCGTAGGCT ACGCTGGCTG CGGCGTGGCG GTGACCTTCT TCCTGTACTT CCTGGCTACC AACTA  887
mPTH1  CGCCGCTGCC GCCGTTGGCT ACGCTGGCTG CGGCGTGGGC GTGACCTTCT TCCTGTACTT CCTGGCTACC AACTA  887
hPTH1  ------TGCC ------GGCT ACGCGGGCTG CAGGGTGGCT GTGACCTTCT TCCTTTACTT CCTGGCTACC AACTA  887

dPTH1  CTACTGGATT CTGGTGGAGG GGCTGTACCT GCATAGCTCTC ATCTTCATGG CCTTGTTCTC AGAGAAGAAG TACCT  959
rPTH1  CTACTGGATT CTGGTGGAGG GGCTGTACTT GCACAGCTC ATCTTCATGG CCTTTTTCTC AGAGAAGAAG TATCT  962
mPTH1  CTACTGGATT CTGGTGGAGG GACTGTACTT ACACAGCTC ATCTTCATGG CCTTTTTCTC AGAGAAGAAG TATCT  962
hPTH1  CTACTGGATT CTGGTGGAGG GGCTGTACCT GCACAGCTC ATCTTCATGG CCTTGTTCTC AGAGAAGAAG TACCT  962
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Figure 2 con't

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dPTH1  GTGGGGCTTC ACGGTCTTTG GCTGGGGTCT GCCGGGGTC TTCGTGGCTG TGTGGGTCAG CGTGAGAGGC ACCGT   1034
rPTH1  GTGGGGCTTC ACCATCTTTG GCTGGGGTCT ACCGGCTGTC TTCGTGGCTG TGTGGGTCGG TGTCAGAGCA ACCTT   1037
mPTH1  GTGGGGCTTC ACCATCTTTG GCTGGGGTCT GCCGGCTGTC TTCGTGGCTG TGTGGGTCGG TGTCAGAGCA ACCTT   1037
hPTH1  GTGGGGCTTC ACAGTCTTCG GCTGGGGTCT GCCGGCTGTC TTCGTGGCTG TGTGGGTCAG TGTCAGAGCT ACCGT   1037

dPTH1  GGCCAACACC GGGTGCTGGG ACTTGAGCTC CGGGAACAAG AAGTGGATCA TCCAGGTGCC CATCCTGGGC TCTAT   1109
rPTH1  GGCCAACACT GGGTGCTGGG ATCTGAGCTC CGGGACAAG AAGTGGATCA TCCAGGTGCC CATCCTGGCA TCTGT   1112
mPTH1  GGCCAACACC GGGTGCTGGG ACTGAGCTC TGGGCACAAG AAGTGGATCA TCCAGGTGCC CATCCTGGGC TCTGT   1112
hPTH1  GGCCAACACC GGGTGCTGGG ACTTGAGCTC CGGGAACAAA AAGTGGATCA TCCAGGTGCC CATCCTGGGC TCCAT   1112

dPTH1  TGTGCTCAAC TTCATCTTGT TCATCAACAT CGTCCGGGTG CTGGCCACCA AGCTGCGGGA GACCAATGCC GGCCG   1184
rPTH1  TGTGCTCAAC TTCATCCTTT TTATCAACAT CATCCGGGTG CTTGCCACTA AGCTTCGGGA GACCAATGCG GGCCG   1187
mPTH1  TGTGCTCAAC TTCATCCTCT TTATCAACAT CGTCCGGGTG CTTGCCACCA AGCTGCGGGA GACCAAGGCC GGCCG   1187
hPTH1  TGTGCTCAAC TTCATCCTGT TCATCAATAT CGTCCGGGTG CTGGCCACCA AGCTGCGGGA GACCAAGGCC GGCCG   1187

dPTH1  GTGTGACACG CGGCAGCAGT ACCGGAAGCT GCTCAAATCC ACAGTGGTGC TCATGCGGCT CTTTGGGGTC CACTA   1259
rPTH1  GTGTGACACC AGGCAGCAGT ACCGGAAGCT GCTCAGGGTC ACGTTGGTGC TTGTGCCACT CTTGGGTGTC CACTA   1262
mPTH1  GTGTGACACA CGGCAGCAGT ACCGGAAGCT GCTCAAATCC ACGGTGGTGC TCATGCGGCT CTTTGGGGTC CACTA   1262
hPTH1  GTGTGACACA CGGCAGCAGT ACCGGAAGCT GCTCAAATCC ACGGTGGTGC TCATGCGGCT CTTTGGGGTC CACTA   1262

dPTH1  CATCGTCTTC ATGGCCACGC CGTACACCGA GGTCTCAGGG ACGCTCTGGC AGTCCAGAT GCATTATGAG ATGCT   1334
rPTH1  CACCGTCTTC ATGGCCTTGC CGTACACCGA GGTCTCACGGG ACATTGTGGC AGATCCAGAT GCACTATGAG ATGCT   1337
mPTH1  CACCGTCTTC ATGGCCTTGC CGTACACCGA GGTCTCAGGG ACACTGTGGC AGATCCAGAT GCACTATGAG ATGCT   1337
hPTH1  CATTGTCTTC ATGGCCACAC CATACACCGA GGTCTCAGGG ACGCTCTGGC AGTCCAGAT GCACTATGAG ATGCT   1337

dPTH1  CTTCAACTCC TTCCAGGGAT TTTTTGTGGC CATCATATAC TGTTTCTGCA ATGGCGAGGT ACAGGCGGAG ATCAA   1409
rPTH1  CTTCAACTCC TTCCAGGGAT TTTTTGTGGC CATCATATAC TGTTTCTGCA ATGGTGAGGT GCAGGCAGAG ATTAG   1412
mPTH1  CTTCAACTCC TTCCAGGGAT TTTTTGTGGC AATCATATAC TGTTTCTGCA ATGGCGAGGT ACAGGCTGAG ATCAA   1412
hPTH1  CTTCAACTCC TTCCAGGGAT TTTTTGTGGC AATCATATAC TGTTTCTGCA ATGGCGAGGT ACAGGCGGAG ATCAA   1412

dPTH1  GAAATCCTGG AGCCGCTGGA CACTGGCCCT GGACTTCAAG CGGCAAAGGCC GAAGTGGGAG CAGCAGTTAC AGCTA   1484
rPTH1  GAAGTCATGG AGCCGCTGGA CACTGGCCGTT GGACTTCAAG CGGCAAAGGCC GAAGTGGGAG TAGCAGCTAC AGCTA   1487
mPTH1  GAAGTCTTGG AGCCGCTGGA CACTGGCACT GGACTTCAAG CGAAAGGAC GCAGCGGGAG CAGCAGCTAT AGCTA   1487
hPTH1  GAAATCTTGG AGCCGCTGGA CACTGGCACT GGACTTCAAG CGAAAGGAC GCAGCGGGAG CAGCAGCTAT AGCTA   1487

dPTH1  CGGCCCCGATG GTGTCTCACA CGAGCGTGAC CAAGGTAGGC CCCGGCGCGG GACTTTGGCCT GCCCCTCAGC CCCCG   1559
rPTH1  TGGCCCCAATG GTGTCTCACA CGAGTGTGAC CAATGTGGGC CCCCGTGCAG GACTCAGCCT TCCCCTTAGC CCCCG   1562
mPTH1  TGGCCCCAATG GGTGCACACA CGAGTGTGAC CAATGTGGGC CCCCGTGCGG GACTCGGCCT GCCCCTCAGC CCCCG   1562
hPTH1  CGGCCCCGATG GTGTCCCACA CAAGTGTGAC CAATGTGGGC CCCCGTGTCG GACTCGGCCT GCCCCTCAGC CCCCG   1562

dPTH1  CCTGCTGGCC GCCGCTGCCG CCACCACCAC GCCGCCACCACC AACGGCCACG CCCCGGATCC GGGCCACACC AAGCC   1634
rPTH1  CCT---GGCT ---------- -------GC TGCCACCACC AATGGCCACT CCCCAGGCTGCC TGGCCACGCC AAGCC   1616
mPTH1  CCT---GGCT ---------- -------GC TGCCACCACC AATGGCCACC CCCCAGGCTGCC TGGCCATGCC AAGCC   1616
hPTH1  CCTACTGGCC ---------- -------AC TGCCACCACC AACGGCCACC CTCCAGGCTGCC TGGCCATGCC AAGCC   1619

dPTH1  AGGGGGCCCG ACCCT----- --CCCGG-GCC CACACCACCT GCCAGGCGTG CTCCCAAGGA CGATGGGTTC CTCAA   1700
rPTH1  AGGGCGCTCCA GCCCCATTGAGA CT---GAAAC CCTACCAGTC ACTATGGACGG TTCCCAAGGA CGATGGGTTC CTCAA   1688
mPTH1  GGGCGCTCCA GCCCTGGAGA AC--GAAAC CATACCAGTT ACTATGACAG TTCCCAAGGA CGATGGGTTC CTCAA   1688
hPTH1  AGGGACCCCA GCCCTGGAGA CCCTCGGAGAC CACACCACCT GCCATGGCTG CTCCCAAGGA CGATGGGTTC CTCAA   1694

dPTH1  CGGCTCCTGC TCGGGGCTGG ACGAGGAGGC CTCGGGGCGG GAGCGGCCCTC CCGCCCTGCT GCAGGACGAG TGGGA   1775
rPTH1  CGGCTCCTGC TCAGGGCTGG ATGAGGAGGC CTCTGGGTCT GCGCGGCCAC CTGCATTGTT GCAGGAAGAA TGGGA   1763
mPTH1  TGGCTCCTGC TCGGGGCTGG ACGAGGAGGC CTCTGGGCCT GAGCGGCCAC CTGCCCTGCT ACAGGAAGAG TGGGA   1769
hPTH1  CGGCTCCTGC TCAGGGCTGG ACGAGGAGGC CTCTGGGCCT GCGCGGCCAC CTGCCCTGCT ACAGGAAGAG TGGGA   1769

dPTH1  GACGGTCATG TGA                                                              1788
rPTH1  AACAGTCATG TGA                                                              1776
mPTH1  AACAGTCATG TGA                                                              1776
hPTH1  GACAGTCATG TGA                                                              1782
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Seq. ID No. 3A dPTH1; Seq. ID No. 3B rPTH1; Seq. ID No. 3C mPTH1 & Seq. ID No. 3D hPTH1

Figure 3

dPTH1	MGAVRIARGL ALLLCCPVLS SAYALVDADD VTKEEQIFL LHRAQAQQK RLKEVLQRPA DIMESDKGWA SASTSGKKK EKASGKLYPE SEEDKEVPTG	100
rPTH1	MGAARIAPSL ALLLCCPVLS SAYALVDADD VTKEEQIFL LHRAQAQQK LLKEVLHTAA NIMESDKGWT PASTSGKPRK EKASGKPRK FTYPE SKENKDVPTG	100
mPTH1	MGTARIAPSL ALLLCCPVLS SAYALVDADD VTKEEQIFL LHRAQAQQK LLKEVLHTAA NIMESDKGWT PASTSGKPRK EKAPGKFTYPE SKENKDVPTG	100
hPTH1	MGTARIARGL ALLLCCPVLS SAYALVDADD VTKEEQIFL LHRAQAQQEK RLKEVLQRPA SIMESDKGWT SASTSGKPRK DKASGKLYPE SEEDKEAPTG	100

dPTH1	SRHRGRPCLP EWDHILCWPL GAPGEVVAVP CPDYIYDFNH KGHAYRRCDR NGSWELVPGH NRTWANYSEC MKFLTNETRE REVFDRLGMI YTVGYSVSLA	200
rPTH1	SRHRGRPCLP EWDNIVCWPL GAPGEVVAVP CPDYIYDFNH KGHAYRRCDR NGSWELVPGH NRTWANYSEC LKFMTNETRE REVFDRLGMI YTVGYSNSLA	200
mPTH1	SRRRGRPCLP EWDNIVCWPL GAPGEVVAVP CPDYIYDFNH KGHAYRRCDR NGSWELVPGH NRTWANYSEC LKFMTNETRE REVFDRLGMI YTVGYSNSLA	200
hPTH1	SRYRGRPCLP EWDHILCWPL GAPGEVVAVP CPDYIYDFNH KGHAYRRCDR NGSWELVPGH NRTWANYSEC MKFLTNETRE REVFDRLGMI YTVGYSVSLA	200

dPTH1	SLTVAVLILA YFRRLHCTRN YIHMHLFLSF MLRAVSIFVK DAVLYSGATL DEAERLTEEE LRAIAQAPPP PTAAAGYAG CRVAVTFFLY FLATNYYWIL	299
rPTH1	SLTVAVLILA YFRRLHCTRN YIHMHNFLSF MLRAASIFVK DAVLYSGFTL DEAERLTEEE LHIIAQVPPP PAAAAVGYAG CRVAVTFFLY FLATNYYWIL	300
mPTH1	SLTVAVLILA YFRRLHCTRN YIHMHNFLSF MLRAASIFVK DAVLYSGFTL DEAERLTEEE LHIIAQVPPP PAAAAVGYAG CRVAVTFFLY FLATNYYWIL	300
hPTH1	SLTVAVLILA YFRRLHCTRN YIHMHLFLSF MLRAVSIFVK DAVLYSGATL DEAERLTEEE LRAIAQAPAT AAAGYAG CRVAVTFFLY FLATNYYWIL	300

dPTH1	VEGLYLHSLI FMAFFSEKKY LWGFTIVFGWG LPAVFVAVWV SVRATLANTG CWDLSSGNKK WIIQVPILAS IVLNFILFIN IMRVLATKLR ETNAGRCDTR	399
rPTH1	VEGLYLHSLI FMAFFSEKKY LWGFTIFGWG LPAVFVAVWV GVRATLANTG CWDLSSGHKK WIIQVPILAS MVLNFILFIN IIRVLATKLR ETNAGRCDTR	400
mPTH1	VEGLYLHSLI FMAFFSEKKY LWGFTIFGWG LPAVFVAVWV GVRATLANTG CWDLSSGHKK WIIQVPILAS MVLNFILFIN IIRVLATKLR ETNAGRCDTR	400
hPTH1	VEGLYLHSLI FMAFFSEKKY LWGFTVFGWG LPAVFVAVWV SVRATLANTG CWDLSSGNKK WIIQVPILAS IVLNFILFIN IMRVLATKLR ETNAGRCDTR	400

dPTH1	QQYRKLLKST LVLMPLFGVH YIVFMAPYT EVSGTLWQVQ MHYEMLFNSF QGFFVAIIYC FCNGEVQAEI KKSWSRWTLA LDFKRKARSG SSSYSYGPNV	499
rPTH1	QQYRKLLKST LVLVPLFGVH YTVFMAIQ PYT EVSGTLWQIQ MHYEMLFNSF QGFFVAIIYC FCNGEVQAEI RKSWSRWTLA LDFKRKARSG SSSYSYGPNV	500
mPTH1	QQYRKLLRST LVLVPLFGVH YIVFMAIQ PYT EVSGTLWQIQ MHYEMLFNSF QGFFVAIIYC FCNGEVQAEI RKSWSRWTLA LDFKRKARSG SSSYSYGPNG	500
hPTH1	QQYRKLLKST LVLMPLFGVH YIVFMAPYT EVSGTLWQVQ MHYEMLFNSF QGFFVAIIYC FCNGEVQAEI KKSWSRWTLA LDFKRKARSG SSSYSYGPNV	500

dPTH1	SHTSVTNVGP RAGLGLPLSP RLLPAAAATT TATTNGHPPI PGHTKPGART L---PATPPA TAAPKDDGFL NGSCSGLDEE ASAPERPPAL LQEEWETVM	595
rPTH1	SHTSVTNVGP RAGLSLPLSP RLPPAT---- TT-NGHSQL PGHAKPGAPA TET-ETLPVT MAVPKDDGFL NGSCSGLDEE ASGSARPPPL LQEEWETVM	591
mPTH1	AHTSVTNVGP RAGLSLPLSP RLLPAT---- TT-NGHSQL PGHAKPGAPA IEN-ETIPVT MTVPKDDGFL NGSCSGLDEE ASGSARPPPL LQEEWETVM	591
hPTH1	SHTSVTNVGP RMGLGLPLSP RLLPTA---- TTNGHPQL PGHAKPGTPA LETLETTPPA MAAPKDDGFL NGSCSGLDEE ASGPERPPAL LQEEWETVM	593

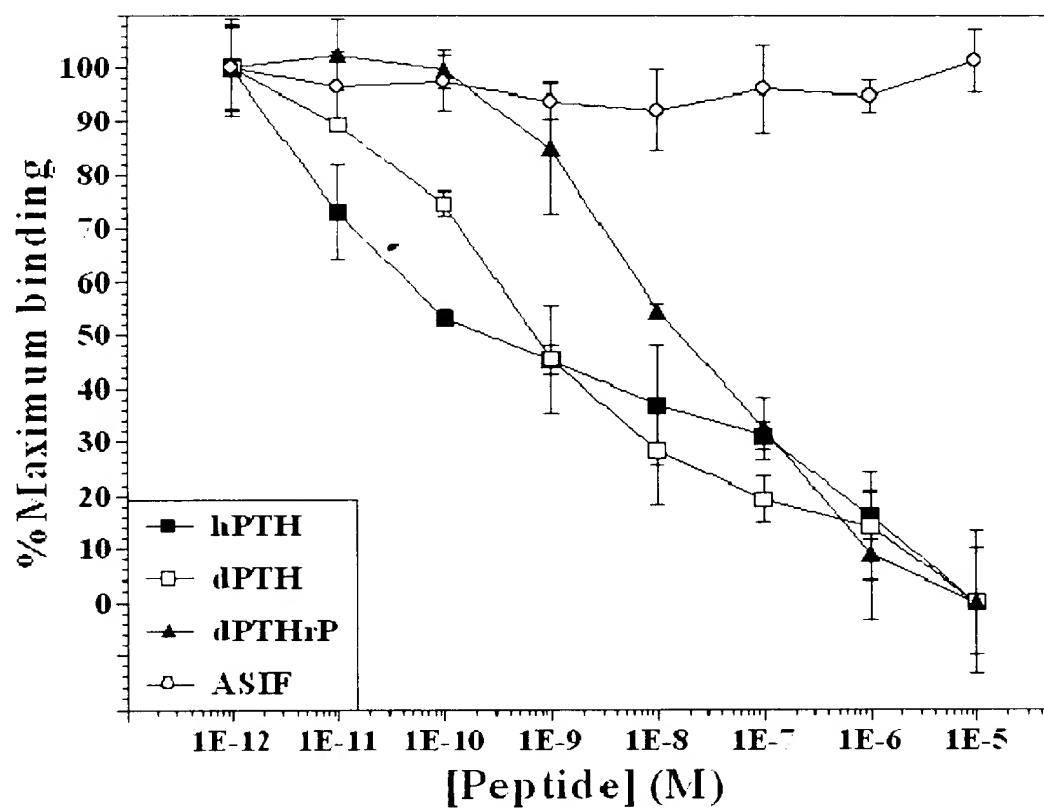Seq. ID No. 2A dPTH1; Seq. ID No. 2B rPTH1; Seq. ID No. 2C mPTH1 and Seq. ID No. 2D hPTH1.

Figure 4

Figure 5